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 M G A U R I A P G L A L  
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 L L C C P U L S S A Y A L U D A D D U M T K E E Q I F L L H A A Q  
 GCGCAGTGCC AGAGCGCGCT CAAAGAGTC CTGCGAGGC CAGCTGACAT AATGGATCA GACAAAGGAT GGGCTTCTGC ATCCACATCA GGAAGCCTA 500  
 A Q C Q K R L K E U L Q R P A D I M E S D K G W A S A S T S G K P K  
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 K E K A S G K L Y P E S E E D K E U P T G S R H A G R P C L P E W  
 GGACCAATC CTCTGCTGGC CGCTGGGGG ACCAGGTGAG GTGGTGGCTG TGCCCTGTCC CCACTACATT TATGACTTCA ATCACAAGG CCACTGCTAC 700  
 D H I L C W P L G A P G E U U A U P C P D Y I Y D F N H K G H A Y  
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 R A C D R N G S W E L U P G H N R T W A N Y S E C U K F L T N E T R  
 GTGACGGGA GGTGTTTGC CGCTGGGA TGATCTACG CAGGGGCTAC TCCGTGTCC TGCCCTCCCT CACCGTGGCC GTGCTCATCC TGCCCTACTT 900  
 E R E U F D R L G H I Y T U G V S U S L A S L T U A U L I L A Y F  
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 R A L H C T R N Y I H M H L F L S F M L R A U S I F U K D A U L Y  
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 S G A T L D E A E R L T E E E L R A I A Q A P P P P T A A A G Y A G  
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 I U R U L A T K L R E T N A G R C D T R Q Q Y R K L L K S T L U L  
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 M P L F G U H V I U F M A T P Y T E U S G T L W Q U Q M H Y E M L  
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 P E R P P A L L Q E E W E T U N  
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Fig. 1

SEQ ID NO: 1



[illegible]

3.

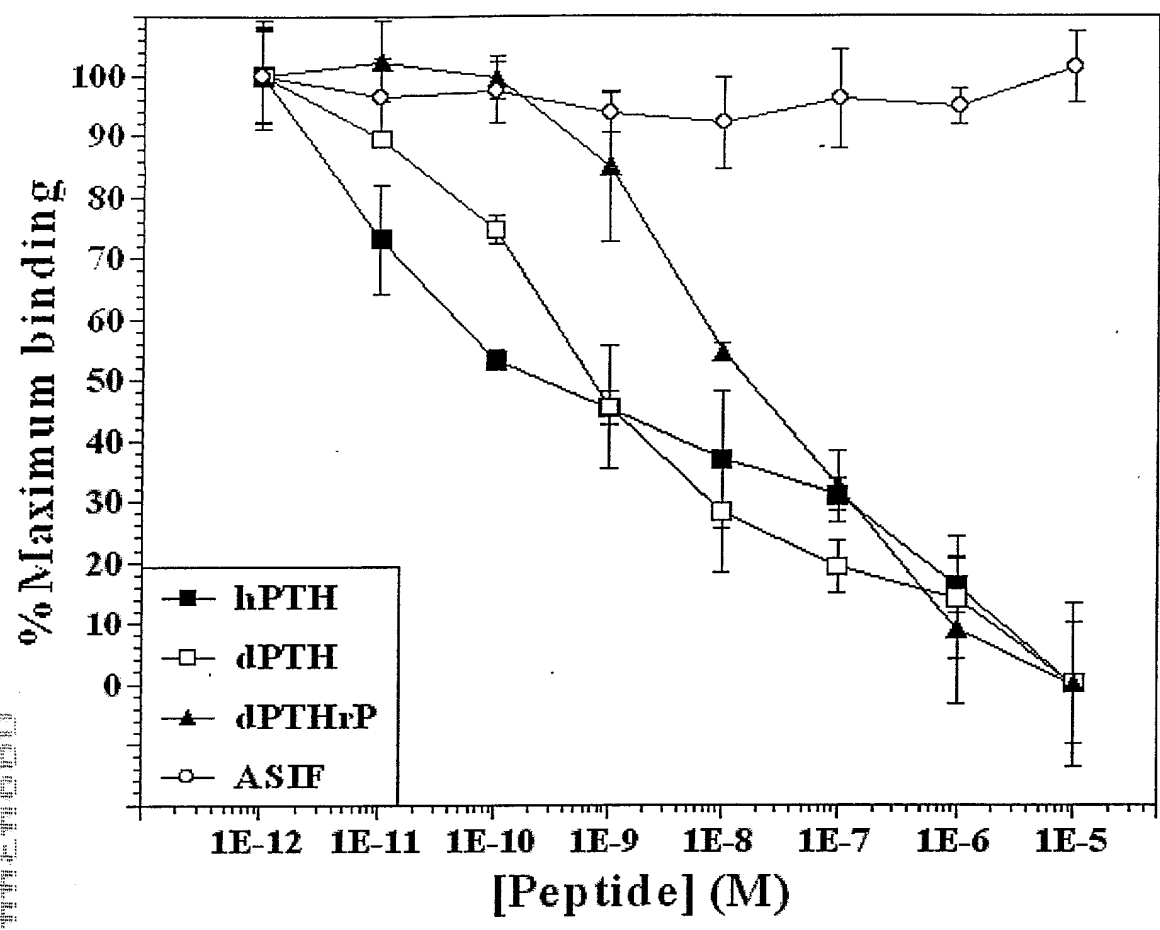


Fig. 4

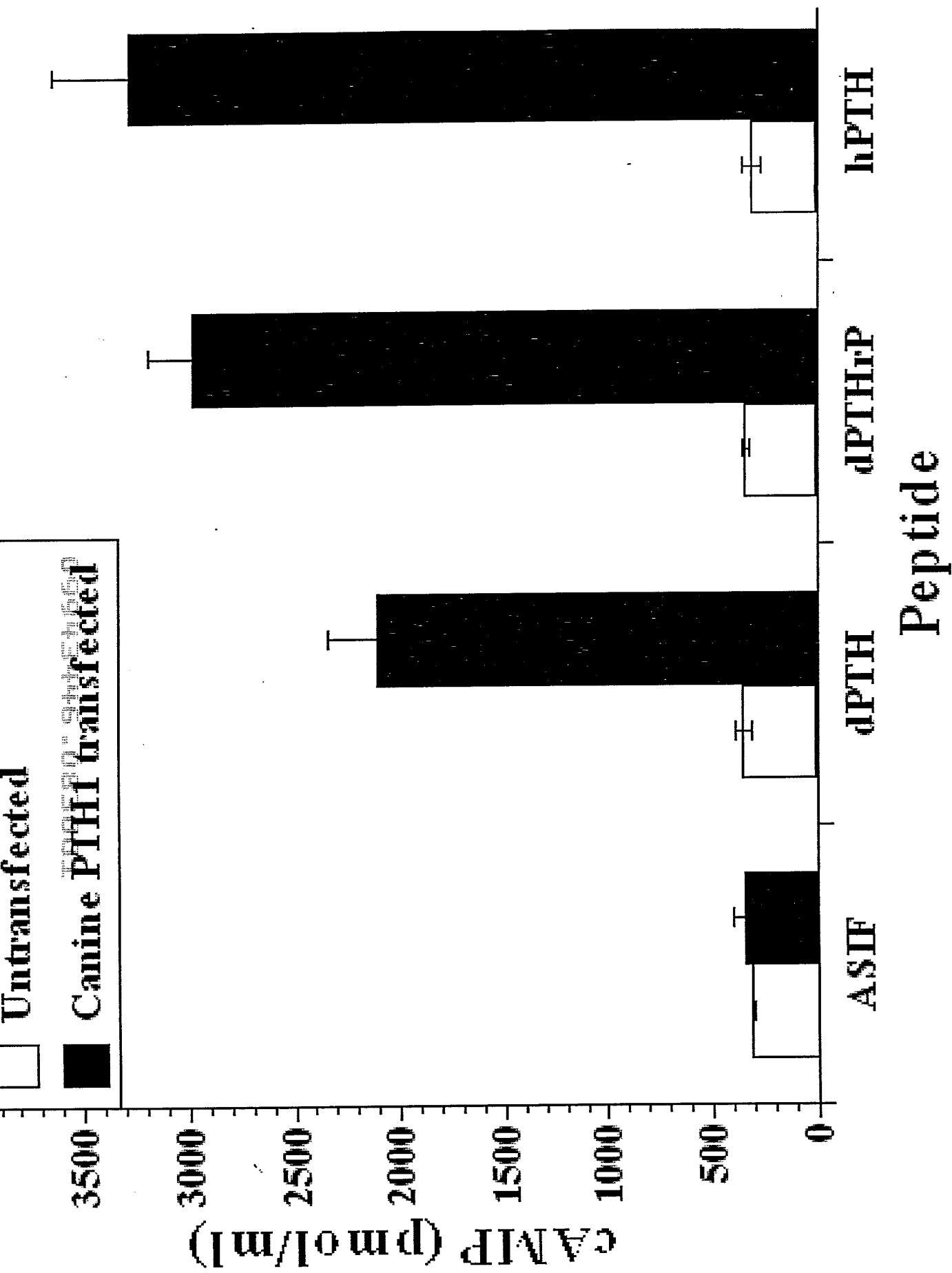


Fig. 5